

## STIC Biotechnology Systems Branch

### RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/522,106

Source:

PCT

Date Processed by STIC:

2-2-05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):  
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



PCT

## RAW SEQUENCE LISTING

DATE: 02/02/2005

PATENT APPLICATION: US/10/522,106

TIME: 15:31:41

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\02022005\J522106.raw

2 <110> APPLICANT: Kogel, Karl-Heinz  
 3 Huckelhoven, Ralph  
 4 Trujillo, Marco  
 6 <120> TITLE OF INVENTION: Method for Obtaining a Pathogen Resistance in Plants  
 8 <130> FILE REFERENCE: 532622010500  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/522,106  
 11 <141> CURRENT FILING DATE: 2005-01-24  
 13 <160> NUMBER OF SEQ ID NOS: 24  
 14 <170> SOFTWARE: PatentIn Ver. 2.1

## ERRORED SEQUENCES

45 <210> SEQ ID NO: 2  
 46 <211> LENGTH: 112  
 47 <212> TYPE: PRT  
 48 <213> ORGANISM: Hordeum vulgare  
 W--> 49 <400> SEQUENCE: 2  
 50 Phe Lys Gly Ile Met Asn Glu Ile Ala Glu Leu Asp Gln Arg Asn Ile  
 51 1 5 10 15  
 52 Ile Glu Met His Asn Tyr Leu Thr Ser Val Tyr Glu Glu Gly Asp Ala  
 53 20 25 30  
 54 Arg Ser Ala Leu Ile Thr Met Leu Gln Ala Leu Asn His Ala Lys Asn  
 55 35 40 45  
 E--> 56 Gly Val Asp Val Val Ser Xaa Thr Arg Val Arg Thr His Phe Ala Arg  
 57 50 55 60  
 58 Pro Asn Phe Lys Arg Val Leu Ser Lys Val Ala Ala Lys His Pro Tyr  
 59 65 70 75 80  
 60 Ala Lys Ile Gly Val Phe Tyr Cys Gly Ala Pro Val Leu Ala Gln Glu  
 61 85 90 95  
 62 Leu Ser Asn Leu Cys His Glu Phe Asn Gly Lys Cys Thr Thr Lys Phe  
 63 100 105 110  
 2181 <210> SEQ ID NO: 16  
 2182 <211> LENGTH: 939  
 2183 <212> TYPE: PRT  
 2184 <213> ORGANISM: Nicotiana tabacum  
 W--> 2185 <400> SEQUENCE: 16  
 2186 Met Gln Asn Ser Glu Asn His His Pro His His Gln His His His Ser  
 2187 1 5 10 15  
 2188 Asp Thr Glu Ile Ile Gly Asn Asp Arg Ala Ser Tyr Ser Gly Pro Leu  
 2189 20 25 30  
 2190 Ser Gly Pro Leu Asn Lys Arg Gly Gly Lys Lys Ser Ala Arg Phe Asn  
 2191 35 40 45

*pls explain*  
*"Xaa" location*  
*Does Not Comply*  
*Corrected Diskette Needed*  
*(pgs. 1, 3)*  
*See error explanation*  
*on page 5.*

## RAW SEQUENCE LISTING

DATE: 02/02/2005

PATENT APPLICATION: US/10/522,106

TIME: 15:31:42

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\02022005\J522106.raw

```

2192 Ile Pro Glu Ser Thr Asp Ile Gly Thr Ser Val Gly Thr Gly Gly Lys
2193      50                      55                      60
2194 Ser Asn Asp Asp Ala Tyr Val Glu Ile Thr Leu Asp Val Arg Glu Asp
2195 65                      70                      75                      80
2196 Ser Val Ala Val His Ser Val Lys Thr Ala Gly Gly Asp Asp Val Glu
2197                      85                      90                      95
2198 Asp Pro Glu Leu Ala Leu Leu Ala Lys Gly Leu Glu Lys Lys Ser Thr
2199                      100                      105                      110
2200 Leu Gly Ser Ser Leu Val Arg Asn Ala Ser Ser Arg Ile Arg Gln Val
2201                      115                      120                      125
2202 Ser Gln Glu Leu Arg Arg Leu Ala Ser Leu Asn Lys Arg Pro Ile Pro
2203 130                      135                      140
2204 Thr Gly Arg Phe Asp Arg Asn Lys Ser Ala Ala Ala His Ala Leu Lys
2205 145                      150                      155                      160
2206 Gly Leu Lys Phe Ile Ser Lys Thr Asp Gly Gly Ala Gly Trp Ala Ala
2207                      165                      170                      175
2208 Val Glu Lys Arg Phe Asp Glu Ile Thr Ala Ser Thr Thr Gly Leu Leu
2209                      180                      185                      190
2210 Pro Arg Ala Lys Phe Gly Glu Cys Ile Gly Met Asn Lys Glu Ser Lys
2211                      195                      200                      205
2212 Glu Phe Ala Val Glu Leu Tyr Asp Ala Leu Ala Arg Arg Arg Asn Ile
2213 210                      215                      220
2214 Thr Thr Asp Ser Ile Asn Lys Ala Gln Leu Lys Glu Phe Trp Asp Gln
2215 225                      230                      235                      240
2216 Val Ala Asp Gln Ser Phe Asp Ser Arg Leu Gln Thr Phe Phe Asp Met
2217                      245                      250                      255
2218 Val Asp Lys Asp Ala Asp Gly Arg Ile Thr Glu Glu Glu Val Arg Glu
2219                      260                      265                      270
2220 Ile Ile Gly Leu Ser Ala Ser Ala Asn Arg Leu Ser Thr Ile Gln Lys
2221                      275                      280                      285
2222 Gln Ala Asp Glu Tyr Ala Ala Met Ile Met Glu Glu Leu Asp Pro Asn
2223 290                      295                      300
2224 Asn Leu Gly Tyr Ile Met Ile Glu Asn Leu Glu Met Leu Leu Leu Gln
2225 305                      310                      315                      320
2226 Ala Pro Asn Gln Ser Val Gln Arg Gly Gly Glu Ser Arg Asn Leu Ser
2227                      325                      330                      335
2228 Gln Met Leu Ser Gln Lys Leu Lys His Thr Gln Glu Arg Asn Pro Ile
2229                      340                      345                      350
2230 Val Arg Trp Tyr Lys Ser Phe Met Tyr Phe Leu Leu Asp Asn Trp Gln
2231                      355                      360                      365
2232 Arg Val Trp Val Leu Leu Leu Trp Ile Gly Ile Met Ala Gly Leu Phe
2233 370                      375                      380
2234 Thr Trp Lys Tyr Ile Gln Tyr Lys Glu Lys Ala Ala Tyr Lys Val Met
2235 385                      390                      395                      400
2236 Gly Pro Cys Val Cys Phe Ala Lys Gly Ala Ala Glu Thr Leu Lys Leu
2237                      405                      410                      415
2238 Asn Met Ala Ile Ile Leu Phe Pro Val Cys Arg Asn Thr Ile Thr Trp
2239                      420                      425                      430
2240 Leu Arg Asn Lys Thr Arg Leu Gly Ala Ala Val Pro Phe Asp Asp Asn

```

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DATE: 02/02/2005

PATENT APPLICATION: US/10/522,106

TIME: 15:31:42

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\02022005\J522106.raw

2241            435                            440                            445  
 2242 Leu Asn Phe His Lys Val Ile Ala Val Ala Ile Ala Leu Gly Val Gly  
 2243            450                            455                            460  
 2244 Ile His Gly Leu Ser His Leu Thr Cys Asp Phe Pro Arg Leu Leu Asn  
 2245 465                            470                            475                            480  
 2246 Ala Ser Glu Glu Glu Tyr Glu Pro Met Lys Tyr Tyr Phe Gly Asp Gln  
 2247                            485                            490                            495  
 2248 Pro Glu Ser Tyr Trp Trp Phe Ile Lys Gly Val Glu Gly Val Thr Gly  
 2249                            500                            505                            510  
 2250 Ile Ile Met Val Val Leu Met Ala Ile Ala Phe Thr Leu Ala Thr Pro  
 2251                            515                            520                            525  
 2252 Trp Phe Arg Arg Asn Arg Val Ser Leu Pro Lys Pro Phe His Lys Leu  
 2253                            530                            535                            540  
 E--> 2254 Thr Gly **Xaa** Asn Ala Phe Trp Tyr Ser His His Leu Phe Val Ile Val  
 2255 545                            550                            555                            560  
 2256 Tyr Thr Leu Phe Ile Val His Gly Glu Lys Leu Tyr Ile Thr Lys Asp  
 2257                            565                            570                            575  
 2258 Trp Tyr Lys Arg Thr Asp Met Asp Val Leu Leu Thr Ile Pro Ile Ile  
 2259                            580                            585                            590  
 2260 Leu Tyr Ala Ser Glu Arg Leu Ile Arg Ala Phe Arg Ser Ser Ile Lys  
 2261                            595                            600                            605  
 2262 Ala Val Lys Ile Leu Lys Val Ala Val Tyr Pro Gly Asn Val Leu Ala  
 2263                            610                            615                            620  
 2264 Leu His Met Ser Lys Pro Gln Gly Tyr Lys Tyr Lys Ser Gly Gln Tyr  
 2265 625                            630                            635                            640  
 2266 Met Phe Val Asn Cys Ala Ala Val Ser Pro Phe Glu Trp His Pro Phe  
 2267                            645                            650                            655  
 2268 Ser Ile Thr Ser Ala Pro Gly Asp Asp Tyr Leu Ser Val His Ile Arg  
 2269                            660                            665                            670  
 2270 Thr Leu Gly Asp Trp Thr Arg Gln Leu Lys Thr Val Phe Ser Glu Val  
 2271                            675                            680                            685  
 2272 Cys Gln Pro Pro Pro Asn Gly Lys Ser Gly Leu Leu Arg Ala Asp Tyr  
 2273                            690                            695                            700  
 2274 Leu Gln Gly Glu Asn Asn Pro Asn Phe Pro Arg Val Leu Ile Asp Gly  
 2275 705                            710                            715                            720  
 2276 Pro Tyr Gly Ala Pro Ala Gln Asp Tyr Lys Lys Tyr Glu Val Val Leu  
 2277                            725                            730                            735  
 2278 Leu Val Gly Leu Gly Ile Gly Ala Thr Pro Met Ile Ser Ile Val Lys  
 2279                            740                            745                            750  
 2280 Asp Ile Val Asn Asn Met Lys Ala Met Asp Glu Glu Glu Asn Ser Leu  
 2281                            755                            760                            765  
 2282 Glu Asp Gly His Asn Asn Asn Met Ala Pro Asn Ser Ser Pro Asn Ile  
 2283                            770                            775                            780  
 2284 Ala Lys Asn Lys Gly Asn Lys Ser Gly Ser Ala Ser Gly Gly Asn Asn  
 2285 785                            790                            795                            800  
 2286 Phe Asn Thr Arg Arg Ala Tyr Phe Tyr Trp Val Thr Arg Glu Gln Gly  
 2287                            805                            810                            815  
 2288 Ser Phe Asp Trp Phe Lys Gly Ile Met Asn Glu Ala Ala Glu Met Asp  
 2289                            820                            825                            830

↖  
 p/s  
 explain  
 Xaa location.

p/s see  
 error  
 explanation  
 on page 5.

The type of errors shown exist throughout  
 the Sequence Listing. Please check subsequent  
 sequences for similar errors.

## RAW SEQUENCE LISTING

DATE: 02/02/2005

PATENT APPLICATION: US/10/522,106

TIME: 15:31:42

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\02022005\J522106.raw

```
2290 His Lys Gly Val Ile Glu Met His Asn Tyr Cys Thr Ser Val Tyr Glu
2291      835      840      845
2292 Glu Gly Asp Ala Arg Ser Ala Leu Ile Thr Met Leu Gln Ser Leu His
2293      850      855      860
2294 His Ala Lys Asn Gly Val Asp Ile Val Ser Gly Thr Arg Val Lys Ser
2295 865      870      875      880
2296 His Phe Ala Lys Pro Asn Trp Arg Asn Val Tyr Lys Arg Ile Ala Leu
2297      885      890      895
2298 Asn His Pro Glu Ala Lys Val Gly Val Phe Tyr Cys Gly Ala Pro Ala
2299      900      905      910
2300 Leu Thr Lys Glu Leu Arg Gln His Ala Leu Asp Phe Ser His Lys Thr
2301      915      920      925
2302 Ser Thr Lys Phe Asp Phe His Lys Glu Asn Phe
2303      930      935
```

## VARIABLE LOCATION SUMMARY

DATE: 02/02/2005

PATENT APPLICATION: US/10/522,106

TIME: 15:31:43

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\02022005\J522106.raw,

Use of n's or Xaa's (NEW RULES):Use of n's and/or Xaa's have been detected in the Sequence Listing.Use of <220> to <223> is MANDATORY if n's or Xaa's are present.in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:1; Xaa Pos. 55

Seq#:2; Xaa Pos. 55

Seq#:15; N Pos. 1952

Seq#:15; Xaa Pos. 547

Seq#:16; Xaa Pos. 547

## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/522,106

DATE: 02/02/2005

TIME: 15:31:43

Input Set : A:\Sequence Listing.txt

Output Set: N:\CRF4\02022005\J522106.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:15 M:283 W: Missing Blank Line separator, <210> field identifier  
L:19 M:283 W: Missing Blank Line separator, <220> field identifier  
L:23 M:283 W: Missing Blank Line separator, <400> field identifier  
L:34 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:193  
L:49 M:283 W: Missing Blank Line separator, <400> field identifier  
L:56 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:2  
L:69 M:283 W: Missing Blank Line separator, <220> field identifier  
L:73 M:283 W: Missing Blank Line separator, <400> field identifier  
L:255 M:283 W: Missing Blank Line separator, <400> field identifier  
L:379 M:283 W: Missing Blank Line separator, <220> field identifier  
L:383 M:283 W: Missing Blank Line separator, <400> field identifier  
L:570 M:283 W: Missing Blank Line separator, <400> field identifier  
L:698 M:283 W: Missing Blank Line separator, <220> field identifier  
L:702 M:283 W: Missing Blank Line separator, <400> field identifier  
L:904 M:283 W: Missing Blank Line separator, <400> field identifier  
L:1032 M:283 W: Missing Blank Line separator, <220> field identifier  
L:1036 M:283 W: Missing Blank Line separator, <400> field identifier  
L:1234 M:283 W: Missing Blank Line separator, <400> field identifier  
L:1364 M:283 W: Missing Blank Line separator, <220> field identifier  
L:1368 M:283 W: Missing Blank Line separator, <400> field identifier  
L:1556 M:283 W: Missing Blank Line separator, <400> field identifier  
L:1680 M:283 W: Missing Blank Line separator, <220> field identifier  
L:1684 M:283 W: Missing Blank Line separator, <400> field identifier  
L:1868 M:283 W: Missing Blank Line separator, <400> field identifier  
L:1990 M:283 W: Missing Blank Line separator, <220> field identifier  
L:1994 M:283 W: Missing Blank Line separator, <400> field identifier  
L:2102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:1935  
M:341 Repeated in SeqNo=15  
L:2185 M:283 W: Missing Blank Line separator, <400> field identifier  
L:2254 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:16  
L:2309 M:283 W: Missing Blank Line separator, <220> field identifier  
L:2313 M:283 W: Missing Blank Line separator, <400> field identifier  
L:2477 M:283 W: Missing Blank Line separator, <400> field identifier  
L:2589 M:283 W: Missing Blank Line separator, <220> field identifier  
L:2593 M:283 W: Missing Blank Line separator, <400> field identifier  
L:2763 M:283 W: Missing Blank Line separator, <400> field identifier  
L:2879 M:283 W: Missing Blank Line separator, <220> field identifier  
L:2883 M:283 W: Missing Blank Line separator, <400> field identifier  
L:3059 M:283 W: Missing Blank Line separator, <400> field identifier  
L:3179 M:283 W: Missing Blank Line separator, <220> field identifier  
L:3182 M:283 W: Missing Blank Line separator, <400> field identifier  
L:3190 M:283 W: Missing Blank Line separator, <220> field identifier  
L:3193 M:283 W: Missing Blank Line separator, <400> field identifier